Amendment to the Specification

Please replace 3rd paragraph on page 4 with the following paragraph:

"Fig. 4 are sequences of pMLP1 attP (region containing the attP site) (SEQ ID NO: 10), M. carbonacea attB region (SEQ ID NO: 4), pSPRH840 attB/attP left juncture region(SEQ ID NO: 5), pSPRH840 attP/attB right juncture region(SEQ ID NO: 6), (labeled add, respectively). Regions of attP, attB, attB/attP sharing homology are indicated by bold larger sized font. 4a, pSPRH840 attP site indicated by large bold font; 4b, M. carbonacea attB site indicated by large bold font. Arrows indicate: a, attP (pSPRH840 attachment site); b, M. carbonacea tRNA-His gene and attB site; c, pSPRH840 attB/attP tRNA-His gene; d, pSPRH840 attP/attB 3' region of tRNA-His gene. Inverted repeats are indicated by small arrows."

Please replace 4th paragraph on page 4 with the following paragraph:

"Fig. 5 are sequences of pMLP1 attP (region containing the attP site) (SEQ ID NO: 10), M. halophitica attB region (SEQ ID NO: 7), pSPRH840 attB/attP left juncture region (SEQ ID NO: 8), pSPRH840 attP/attB right juncture region (labeled a-d, respectively) (SEQ ID NO: 9). Regions of attP, attB, attB/attP and attP/attB sharing homology are indicated by bold larger sized font. 5a, pSPRH840 attP site indicated by large bold font; 5b, M. halophitica attB site indicated by large bold font. Arrows indicated: a, attP (pSPRH840 attachment site); b, M. halophitica tRNA-His gene and attB site; c, pSPRH840 attB/attP tRNA-His gene; d, pSPRH840 attP/attB 3' region of tRNA-His gene. k Inverted repeats are indicated by small arrows."

Remarks:

Claims 1, 6, 8-18, 21 and 24-26 are pending. New claims 24-26 has been added to the application. Written support for new claim 24 appears in the specification, for example, in the Sequence Listing and at page 3, lines 3-6. Written support for new claims 25 and 26 appear in the specification for example, at page 3, lines 20-27; page 8, lines 21-27; and at page 20, line 23 to page 23, line 13. Only formal changes are made to the other claims that do not add any new matter to the present application.

Election/Restriction. Applicants acknowledge the Examiner's invitation to rejoin SEQ ID NO: 6 into the claims of the elected group. SEQ ID NO: 6 is claimed in new claim 24.

Amendment Practice. Applicants wish to thank the Examiner for agreeing to proceed with examination in spite of the non-compliancy of the May 10, 2004 amendment. Future amendments will be in compliance with 37 C.F.R. § 1.121.

Sequence Compliance. The Examiner has objected to the specification, stating that the sequences appearing in the specification, at pages 13, 23 and in the figures, must be identified by a sequence identifier (SEQ ID NO:).

Pages 13 and 23, in the specification, were amended in the "Preliminary Amendment and Response to Notice to Comply with Sequence Rules" filed on July 16, 2002 to insert sequence identifiers next to the sequences.

The "Brief Description of the Drawings" section, on page 4, was amended herein to insert sequence identifiers where appropriate.

The Examiner also alleges that no sequence listing (paper or computer readable format) or sequence listing statement was ever filed in the present application. Applicants respectfully disagree. The Examiner's attention is directed to the enclosed copy of a postcard bearing a USPTO sticker acknowledging

receipt of a "sequence listing statement", a "sequence disk" and a "sequence listing paper copy" in the present application (Exhibit B). A substitute sequence listing (paper and computer readable format) and statement was submitted on July 16, 2002. In any event, to expedite prosecution of the present application, applicants herein resubmit these items.

Withdrawal of the objection is requested.

Claim Rejections Under 35 U.S.C. § 112. Claim 3 stands rejected because there is allegedly no antecedent support for "said polypeptide". The rejection of claim 3 is moot because the claim has been cancelled.

Claim Rejections Under 35 U.S.C. § 102(e). Claims 1-4, 6, 8-18 and 20-21 stand rejected as anticipated by Hosted et al. (US 2004/0101832). The rejection of claims 2-4 and 20 are moot since the claims have been cancelled. The Examiner alleges the claims are anticipated because both Hosted et al. and the instant claims disclose nucleic acids comprising an M.carbonacea-specific integrase gene and the cognate integration site.

Applicants submit that the rejection should be withdrawn because the claims, as amended, are not anticipated by Hosted et al. Moreover, rejection of new claims 25 and 26 over Hosted et al. is not warranted because the recited method is patentable.

The rejection of claim 1, as it relates to SEQ ID NOs: 1 and 3, is most since the amended claim does not recite these sequences.

Moreover, the Examiner does not appear to allege that claimed SEQ ID NO: 2 (Xis gene) is anticipated by Hosted et al. The Examiner's only alleges that Hosted et al. discloses the claimed "integrase gene" and the "integration sites".

Moreover, it is unclear where, if anywhere, in the cited Hosted et al. reference, that SEQ ID NO: 2 is disclosed. SEQ ID NO: 2

bears no significant sequence homology to SEQ ID NO: 176 of the Hosted et al. reference. As proof of the lack of sequence homology, enclosed is a BlastN output comparing the sequences (Exhibit A). In the BlastN output, the "query" sequence is SEQ ID NO: 176 of the Hosted et al. reference and the "subject" sequence is SEQ ID NO: 2 of the instant application.

Hosted et al. does not anticipate amended claim 6 because, as stated above, the reference does not disclose a polynucleotide or vector comprising the nucleotide sequence of SEQ ID NO: 2.

Hosted et al. does not anticipate claims 8-18 and 21, as amended, or new claims 25 or 26 because the claims specify that the vector comprises SEQ ID NO: 2. In contrast, as discussed above, Hosted et al. does not disclose a polynucleotide comprising the nucleotide sequence of SEQ ID NO: 2.

Conclusion:

Applicants submit that the amended claims are in condition for passage to allowance. Such action is earnestly solicited.

Respectfully submitted,

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